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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.
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OM nucleic - nucleic search, using sw model Title: Perfect score: Run on: Scoring table: Sequence: US-09-675-650-2 278 1 caggaagcacaaaa IDENTITY_NUC Gapop 10.0 , Gapext 1.0 February 1, 2004, 08:17:56; Search time 1513 Seconds (without alignments) 7516.772 Million cell updates/sec caggaagcacaaaaggaagc.....ggcttcacaagacatgcaac 278

Minimum DB seq length: 0
Maximum DB seq length: 200000000 Total number of hits satisfying chosen parameters: 5777422

2888711 seqs, 20454813386 residues

Searched:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : GenEmbl:*
1: gb_ba:*
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	N	N	N	N	c 21	N	щ	ш	_	_	_	_	<u>, , , , , , , , , , , , , , , , , , , </u>	, <u></u>	11:	10			1 0						,	No.	Result	
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ALIGNMENTS

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JOURNAL	TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AX107858	RESULT 1
Patent: WO 0123550-A 2 05-APR-2001;	Pca3 messenger rna species in benign and malignant prostate tissues	Busse, U., Chypre, C. and Fradet, Y.	1	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	Homo sapiens (human)	•	AX107858.1 GI:13923250	AX107858	Sequence 2 from Patent WO0123550.	AX107858 278 bp DNA linear PAT 30-APR-2001		

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Patent: US 6321716-A 471 27-NOV-2001;
Location/Qualifiers
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Mashiki, Z. and Harada, J.
                                                                                                                                                                                                                                                                                              Unknown.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 2e-68;
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Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y., Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H., Vedwick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A.W., Hepler,W.T. and Henderson,R.A.

Compositions and methods for the therapy and diagnosis of p
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Unknown.
            Skeiky,Y.A., Xu,J., Cheever,M.A. and Reed,S.G. Compositions and methods for wtl specific immunotherapy Patent: Wo 0125273-A 345 12-APR-2001; CORIXA CORPORATION (US)
                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                             Sequence 345 from Patent AX106207
                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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189 c 169 g
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Best Local Similarity
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Best Local (
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                                                                                                                                               1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
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Sequence 471 from Patent
AX106690
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                 Similarity
                                                                            TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                  CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
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AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
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CORPORATION (US)
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
189 c 169 g 25
                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
189 c 169 g 25
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99.6%;
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99.6%;
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Pred. No. 2e-68;
0; Mismatches
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Pred. No. 2e-68;
0; Mismatches
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 471 from Patent
AX140981
AX140981.1 GI:14281078
Хu, J.,
                                                                                      Sequence 471 from Paten AX200841 AX200841.1 GI:15390744
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                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent: WO 0134802-A 471 17-MAY-2001; CORIXA CORPORATION (US)
                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer
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 Dillon, D.C.,
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
189 c 169 g 25
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99.6%;
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 Mitcham, J.L., Harlocker, S.L., Jiang, Y.,
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Pred. No. 2e-68;
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WO0134802.
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Day,C.H.,
                                       Euteleostomi;
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REFERENCE
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Matches 277;
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Best Local Similarity
Matches 277; Conser
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Sequence 471 from Patent
AX267497
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                                                                                                                                                                                                                                                                                                                                                     Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y.,
Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H.,
Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A., Hep
and Henderson,R.A.
                                                                                                                                                                                                                                                                          cancer
Patent: WO 0173032-A 471 04-OCT-2001;
CORIXA CORPORATION (US)
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CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
189 c 169 g 25
                                                                                                                                                        /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
189 c 169 g 256
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PD 13-NOV-2001
PF 09-APR-1998 JP 1998542194
PR 10-APR-1997 US 60/041836
PI MARION J G BUSSEMAKERS
PC C12N15/00,C12N15/12,C12N5/12,C12N1/: C12N15/00,C12N15/12,C12N5/12,C12N1/: C12N15/00,C12N15/12,C12N1/: C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N15/12,C12N1
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JP 2001522240-A/3.
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                                                  AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
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13-NOV-2001
09-APR-1998 JP 1998542194
10-APR-1997 US 60/041836
MARION J G BUSSEMAKERS
C12N15/00,C12N15/12,C12N5/12,C12N1/21,C07K14/47,C07K16/18,
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/mol_type="genomic DNA"
/db_xref="taxon:10509"
169 c 191 g 198
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Pred. No. 2e-68;
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AUTHORS
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ORGANISM
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PN JP 2001522240-A/2
PN JP 2001522240-A/2
PF 10-APR-1997 US 60/041836
PI MARION J G BUSSEMAKERS
PC C12N15/00,C12N15/12,C12N5/1
C12Q1/68,
PC G01N33/577,A61K39/395,A01K6
CC Strandedness: Double;
CC Topology: Linear;
FH Key
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                                             BD082482 2037 PCA3, PCA3 genes, and methods BD082482
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                     BD082482.1 GI:22628092
JP 2001522240-A/1.
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                                                                                                                                             AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
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13-NOV-2001
03-APR-1998 JP 1998542194
10-APR-1997 US 60/041836
MARION J G BUSSEMAKERS
C12N15/00,C12N15/12,C12N5/12,C12N1/21,C07K14/47,C07K16/18,
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Strandedness: Double;
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1. .1872
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/mol_type="genomic DNA"
/db_xref="taxon:10509"
389 c 369 g 539 1
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99.6%;
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Pred. No. 2.2e-68;
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PN JP 2001522240-A1
PF 13-MOV-2001
PF 09-APR-1998 JP 198542194
PR 10-APR-1997 US 60/041836
PI MARION J G BUSSEMAKERS
PC C12N15/00,C12N15/12,C12N5/12,C12N1/21,C0
C12Q1/68,
PC G01N33/577,A61K39/395,A01K67/027
CC Strandedness: Double;
PC Topology: Linear;
FH Key
FT CDS 13/57.351.
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                                                                           Negative pressure control apparatus for Patent: US 6321716-A 469 27-NOV-2001; Cocation/Qualifiers
                                                                                                                                1 (bases 1 to 2229)
Mashiki, Z. and Harada, J.
                                                                                                                                                                     Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Viruses; dsDNA viruses, no RNA stage; Adenoviridae.
1 (bases 1 to 2037)
                                                                                                                                                          Unclassified
                                                                                                                                                                                       Unknown
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10-APR-1997 US 60/041836
MARION J G BUSSEMAKERS
C12N15/00,C12N15/12,C12N5/12,C12N1/21,C07K14/47,C07K16/18,
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/mol_type="genomic DNA"
/db_xref="taxon:10509"
426 c 406 g 575 t
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                                                   /organism="unknown"
447 c 481 g
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Pred. No. 2.2e-68;
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RESULT 13
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Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y. Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H. Vedwick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A.W., Hepler,W.T. and Henderson,R.A.

Compositions and methods for the therapy and diagnosis of
                                                                                                                                                                                                                                                                                                                                          Patent: US 6512094-A 469 28-JAN-2003;
Location/Qualifiers
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AR278568
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unclassified.
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                                                             AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                              CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
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RESULT 14

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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           /organism="Homo sapiens"
/mol_type="genomic DNA"
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Db 1378 AGACGAGAAAATC Search completed: February Job time : 1515 secs	181 1437	Qy 121 AATTGJ Db 1497 AATTGJ	Qy 61 TCATCO	Qy 1 CAGGAI Db 1617 CAGGAI	Query Match Best Local Similarity Matches 277; Conser
TTGATGGCTTCACAAGACATGCAAC 1, 2004, 10:18:49	AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC 240	AATTGATGTGTTCCTTAAAGGATGGGCAGGAAAACAGATCCTGTTGTGGATATTTATT	TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG 120 	CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG 60	stch 95.7%; Score 266; DB 6; Length 2229; sal Similarity 99.6%; Pred. No. 2.2e-68; 277; Conservative 0; Mismatches 0; Indels 1; Gaps 1;

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Best Local Similarity 99.6%;
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APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xi Jiangchun
APPLICANT: Xi Jiangchun
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                         APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Lou
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Retter, Mark
Solk, John
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Reed, Steven G.
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APPLICANT: Xu, Jiangcnun
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIA
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR
FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 469
LENGTH: 2229
TYPE: DNA
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; Sequence 469, Application US/09352616A
; Patent No. 6395278
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APPLICANT: Dillon, Dav
APPLICANT: Harlocker,
APPLICANT: Jiang, Yuqu
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Best Local Similarity
Matches 277; Conserv
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 469
LENGTH: 2229
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                                                                                                                                                                                                                                     Best Local Similarity
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Xu, Jiangchun
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Pred. No. 1.6e-79;
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Pred. No. 1.6e-79;
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APPLICANT: Day, Craig
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REPERENCE: 210121 427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOPTWARE: FASKSEQ for Windows Version 3.0
SEQ ID NO 470
LENGTH: 2426
TYPE: DNA
ORGANISM: Homo sapiens
US-09-439-313-470
                                                                                                                        RESULT 6
US-09-52-616A-470/c
; Sequence 470, Application
; Patent No. 6395278
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US-09-439-313-470/c
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                                                      GENERAL INFORMATION:
APPLICANT: Dillon, I
APPLICANT: Harlocker
APPLICANT: Jiang, Yu
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Best Local Similarity
Matches 277; Conserv
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                      APPLICANT:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
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Reed, Steven G.
Kalos, Michael
Fanger, Gary
Retter, Mark
Solk, John
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FILE REFERENCE: 210121.427C8
FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 470
LENGTH: 2426
TYPE: DNA
ORGANISM: Homo sapiens
US-09-352-616A-470
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                                                                                                                                       ; ORGANISM: Homo sapiens US-09-439-313-468
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                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 468
LENGTH: 3112
                                                                Query Match
Best Local Similarity
Matches 277; Conserv
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Best Local Similarity
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
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Mitcham, Jennifer L.
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Pred. No. 1.6e-79;
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                                                                                    Score 266; DB 4;
Pred. No. 1.8e-79;
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; SOFTWARE: FastSEQ for Wi
; SEQ ID NO 468
; LENGTH: 3112
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-352-616A-468
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; Sequence 468, Application US/09352616A
; Patent No. 6395278
                                                                                            RESULT 9
US-09-439-313-313
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Sequence 313, Application US/09439313
PATENT NO. 6329505
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
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Best Local Similarity
Matches 277; Conserv
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APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
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99.6%;
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APPLICANT: Fanger, Gary
APPLICANT: Retter, Mark
APPLICANT: Solk, John
APPLICANT: Bay, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 313
LENGTH: 718
TYPE: DNA
OPCANIEM: Homo sapien
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                                           CURRENT FILING DATE: 1999-07-13

CURRENT FILING DATE: 1999-07-13

NUMBER OF SEQ ID NOS: 472

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 313

LENGTH: 718
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 98.6
Matches 274; Conservative
                                                                                                                                                          APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR L
FILE REFERENCE: 210121.427C8
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               TYPE: DNA ORGANISM: Homo sapien
FEATURE:
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98.6%;
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APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C6
CURRENT APPLICATION NUMBER: US/09/232,149A
CURRENT FILING DATE: 1999-01-15
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 313
LENGTH: 718
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Best Local Similarity
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NAME/KEY: misc feature

LOCATION: (1)...(718)

OTHER INFORMATION: n = A,T,C or

-09-232-149A-313
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LOCATION: (1)...(718)
OTHER INFORMATION: n = A,T,C
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Local Similarity 98.6%;
les 274; Conservative
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                                                          AATTGATGTGTTCCTTAAAGGAT-GGCAGGAAAACAGATCCTGTTGTGGATATTTATTTG 409
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APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Minng, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIA
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR
FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
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                                                  SOFTWARE: FASTSEQ for V
SEQ ID NO 287
LENGTH: 301
TYPE: DNA
ORGANISM: Homo sapien
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                                    US-09-352-616A-287
                                                                                                                                                                                                                                                                                                                                                               Sequence 287, Application US/09352616A Patent No. 6395278 GENERAL INFORMATION:
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SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 287
LENGTH: 301
TYPE: DNA
ORGANISM: Homo sapien
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Patent No. 6329505
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Best Local
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CURRENT FILING DATE: 1999-11-12
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
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Day, Craig
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Reed, Steven G
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Retter, Mark
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100.0%; Pred. No.
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4.9e-11;
Length 301;
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Query Match

21.9%;

Score 61;

DB 4

Best Local Similarity

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; ORGANISM: Homo sapien
US-09-232-149A-287
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CURRENT FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 338
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 287
LENGTH: 301
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14, Application US/08232463 Patent No. 5670367
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Sequence 287, Application US/09232149A
Patent No. 6465611
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                                                                      ZIP: 22313-0299

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, V

CURRENT APPLICATION DATA:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C6
                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                       APPLICANT: SCHEIFLINGER, APPLICANT: FALKNER, F. G
APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 5
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CITY: Alexandria
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Search completed: February Job time: 53 secs
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US-08-232-463-14
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SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pair
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NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 3047
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
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APPLICATION NUMBER:
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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278
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Copyright (c) 1993 - 2004 Compugen Ltd.
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6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	200	266	266	266	266	266	266	266	266	266	266	266	266	266	266	Score
	01.	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	Query
	3	2229	2229	2229	2229	876	812	812	812	812	812	812	812	812	812	Length
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US-10-294-025-690	US-10-144-678A-690	US-09-895-814-690	US-09-895-793-690	US-09-822-827-690	US-09-780-669-690	US-09-759-143-690	US-10-010-940-468	US-10-012-896-468	US-10-294-025-468	US-10-144-678A-468	US-09-895-814-468	US-09-895-793-468	US-09-822-827-468	46	US-09-759-143-468	US-10-205-823-448	-10-010	•	US-10-294-025-470	US-10-144-678A-470	US-09-895-814-470	US-09-895-793-470	US-09-822-827-470	US-09-780-669-470	US-09-759-143-470	US-10-010-940-469	US-10-012-896-469	US-10-294-025-469	-10-144
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-759-143-471
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Patent No. US20020022248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
Query Match
Best Local Similarity 99.0
Matches 277; Conservative
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SEQ ID NO 471
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CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
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APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427023
                                                                                                                      LENGTH: 812
                                                                                                                                                         FastSEQ for Windows Version 3.0
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Carter, Darrick
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Kalos, Michael
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Dillon, Davin C.
Mitcham, Jennifer L.
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ick, Thomas S.
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Score 266; DB 9;
Pred. No. 9.7e-76;
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TYPE: DNA
CORGANISM: Homo sapiens
US-09-780-669-471
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SEQ ID NO 471
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SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: McNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS
TITLE OF INVENTION: DIAGNOSIS OF
FILE REFERENCE: 210121.427C24
                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
                                                                                                                                                Jocal Similarity 99.6%;
les 277; Conservarion
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TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAAGGACATTAGAAAAATG
                                                                                            CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGATCTTTGGG
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Hepler, William
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Carter, Darrick
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Vedvick, Thomas S
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Kalos, Michael D.
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Dillon, Davin C.
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Pred. No. 9.7e-76;
                                                                                                                                                        Mismatches
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Viang, Yuqiu
APPLICANT: Viang, Yuqiu
APPLICANT: Retter, Marc W.
APPLICANT: Retter, Marc W.

APPLICANT: APPLICANT: APPLICANT:

Jiang, Yuqiu
Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick

Samuel X.

Sequence 471, Application US/09895793 Publication No. US20020192763A1.

GENERAL INFORMATION:

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; ORGANISM: Homo sapiens
US-09-822-827-471
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US-09-895-793-471/c
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GENERAL INFORMATION:

APPLICANT: XU, JIANGCHUN
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 471
LENGTH: 812
TYPE: NUM
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                          Matches 277;
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                                                                     AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 286
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Pred. No. 9.7e-76;
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Sequence 471, Application U
Publication No. US200201932
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin
APPLICANT: Mitcham, Jenni
APPLICANT: Harlocker, Sus
APPLICANT: Jiang, Yuqiu
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US-09-895-814-471/c
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; LENGTH: 812
; TYPE: DIA
; ORGANISM: Homo sapiens
US-09-895-793-471
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Best Local Similarity
Matches 277; Conserv
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APPLICANT: Foy, Teresa
APPLICANT: Fonger, Gary R.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C2
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SOFTWARE: FastSEQ for Windows Version
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Harlocker, Susan L.
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Henderson, Robert A.
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Hural, John
McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
                                                                 Skeiky, Yasir A.W.
Hepler, William T.
Henderson, Robert A.
                                                                                                                                                 Vedvick, Thomas
Carter, Darrick
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Kalos, Michael D.
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Hepler, William
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Pred. No. 9.7e-76;
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APPLICANT: Fanger, Gary R.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C26
CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 471, Applic
Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
Matches 277; Conserv
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
              APPLICANT: Vinals y de Bassols, Carlota
APPLICANT: Foy, Teresa M.
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Deng, Ta
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C28
CURRENT APPLICATION NUMBER: US/10/144,678A
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Skeiky, Yasir A. W.
Hepler, William T.
Hural, John
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                                                                                                 Houghton, Raymond L. Vinals y de Bassols, Foy, Teresa M.
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; CURRENT FILING DATE: 2002-08-12; NUMBER OF SEQ ID NOS: 1033; SOFTWARE: FastSEQ for Windows Version 3.; SEQ ID NO 471; LENGTH: 812; TYPE: DNA GRGANISM: Homo sapiens
US-10-144-678A-471:
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; ORGANISM: Homo sapiens
US-10-294-025-471
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Best Local S
Matches 277
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CURRENT APPLICATION NUMBER: US/10/294,025
CURRENT FILING DATE: 2002-11-12
CURRENT FILING DATE: 2002-11-12
NUMBER OF SEQ ID NOS: 1038
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 471
LENGTH: 812
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APPLICANT: Stolk, John A.
APPLICANT: Kalos, Michael D.
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Pred. No. 9.7e-76;
0; Mismatches 0;
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin
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NUMBER OF SEQ ID NOS: 1011
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 471
LENGTH: 812
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Publication No.
                                                                                                                                                                                                                                                                                                         Query Match 95.7%;
Best Local Similarity 99.6%;
Matches 277; Conservative
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APPLICANT: Wantanabe, Yoshihiro
APPLICANT: Meagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIACHOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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Mitcham, Jennifer L.
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Vedvick, Thomas S.
Carter, Darrick
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Kalos, Michae
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Hepler, William T.
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o. US20020183251A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Teresa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Susan L.
                                                            TTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAAGGAAAAC
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                                                                                                                                                                                                                                                                                                                         Score 266; DB 14;
Pred. No. 9.7e-76;
                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                         Length 812;
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US-10-010-940-471/c

Sequence 471, Applic. Publication No. US20 GENERAL INFORMATION:

Application US/10010940 o. US20030088062A1

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APPLICANT: Sun, Yongming
APPLICANT: Recipon, Herve
APPLICANT: Recipon, Herve
APPLICANT: Recipon, Herve
APPLICANT: Ali, Sunjath
TITLE OF INVENTION: Compositions and Methods Relating to
TITLE OF INVENTION: Genes
FILE REFERENCE: DEX-0239
CURRENT APPLICATION NUMBER: US/09/957,708
CURRENT FILING DATE: 2001-09-19
PRIOR APPLICATION NUMBER: 60/233,746
PRIOR FILING DATE: 2000-09-19
DRIOR FILING DATE: 2000-09-19
TRIOR FILING DATE: 2000-09-19
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-010-940-471
                                                                                                                                                                                                                                                                                                             US-09-957-708-3
                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  Sequence 3, Application US/09957708 Publication No. US20030031678A1
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Best Local Similarity
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APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427D3
CURRENT APPLICATION NUMBER: US/10/010,940
CURRENT FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 575
NUMBER OF SEQ ID NOS: 575
NUMBER OF SEQ ID NOS: 575
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181 AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
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Retter, Mark
Solk, John
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Reed, Steven G.
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99.6%;
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; SEQ ID NO 3
; LENGTH: 876
; TYPE: DNA
; ORCANISM: Homo sapiens
US-09-957-708-3
                                                                                                                 SOFTWARE: FastSE
SEQ ID NO 469
LENGTH: 2229
LENGTH: 2229
TYPE: DNA
ORGANISM: Homo s
US-09-759-143-469
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                                                Query Match
Best Local Similarity
Matches 277; Conserv
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Best Local Similarity 99.0
Matches 277; Conservative
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                                                                                                                                                                                                        APPLICANT: SkeĬky, Ýasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.437C23
CURRENT FILLING DATE: 2001-01-12
NUMBER: US/09/759,143
CURRENT FILLING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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1617 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
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                 1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCCTGGGAGAAATGCCCCGGCCGCCATCTTGGG
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                                                                                                                                                                                                                                                                                                                                                 FastSEQ for Windows Version
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
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Mitcham, Jennifer L.
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                                                           Conservative
                                                                                                                                  sapiens
                                                                        95.7%;
99.6%;
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99.6%;
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                                                       Score 266; DB 9; L
Pred. No. 1.7e-75;
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CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 469
LENGTH: 2229
TYPE: DNA
ORGANISM: Homo sapiens
US-09-780-669-469
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US-09-780-669-469/c
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                                                                                                                                                                                                                    Matches 277;
                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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APPLICANT: MCNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C24
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AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC 240
                                                            CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
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                                                                                         TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
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Hepler, William
Hural, John
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Vedvick, Thomas S.
Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Harlocker, Susan L.
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Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Henderson, Robert
Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jiang, Yugui
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99.6%;
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                                                                                                                                                                                                                    Score 266; DB 9; Length 2229; Pred. No. 1.7e-75; O; Mismatches O; Indels
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; ORGANISM: Homo sapiens
US-09-822-827-469
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US-09-822-827-469/c
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APPLICANT: XU, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER

FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 469
LENGTH: 2229

LENGTH: 2229
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Best Local Similarity 99.0
Matches 277; Conservative
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WS-09-895-793-469/c

US-09-895-793-469/c

Sequence 469, Application US/09895793

Publication No. US20020192763A1

GENERAL INFORMATION:
APPLICANT: No. Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Jiang, Yuqiu
APPLICANT: Kalos, Michael D.
APPLICANT: Kalos, Michael D.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel X.
APPLICANT: Skelky, Yasir A.W.
APPLICANT: Hepler, William T.
APPLICANT: Henderson, Robert A.

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; ORGANISM: Homo sapiens
US-09-895-793-469
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Best Local Similarity
Matches 277; Conservat
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LENGTH: 2229
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.334C2
CURRENT FAPPLICATION NUMBER: US/09/895,793
CURRENT FILLING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 982
SOFTMARE: FASTSEQ for Windows Version 3.0
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APPLICANT:
APPLICANT:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Jiang, Yuqiu
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T: Fanger, Gary R.
INVENTION: COMPOSITIONS AND METHODS FOR THE
INVENTION: DIAGNOSIS OF PROSTATE CANCER
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Vedvick, Thomas S.
Carter, Darrick
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Hepler, William T.
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                                                    Foy, Teresa
                                                                                                                   Henderson, Robert A.
Hural, John
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o. US20020193296A1
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Pred. No. 1.7e-75;
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; FILE REFERENCE: 210121.427C26
; CURRENT APPLICATION NUMBER: US/09/895,814
; CURRENT FILING DATE: 2001-06-29
; NUMBER OF SEQ ID NOS: 990
; SOFTWARE: FastSEQ for Windows Version 3.0
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 469
; LENGTH: 2229
; TYPE: DNA
; ORGANISM: Homo sapiens
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AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 1341
                       AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                                       AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC 1379
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Pred. No. 1.7e-75;
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Search completed: February Job time : 223 secs 2004, 10:45:36

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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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41.4	43.6	2 119	. 165	esult No. Score
14.9	15.7	42.8	59.4	Query
1016	489	402	167	Query Match Length DB ID
13	10	9	10	DB .
BX442677	BG224987	AA578773	10 BF373406	Query Match Length DB ID
BX442677 BX442677	BG224987 kp52h05.y	AA578773 nh24a04.s	BF373406 IL2-FT015	Description

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AG001111 HOMO BAD1 AZ689940 ENTIMIZTE AZ547772 ENTFQ09TF AG220724 Lotus jap AQ070955 HS 2174 A AQ188863 HS 3218 A BI324783 rk88910.y	BZ797594 PUGDT71TD CC384522 PUHBM10TD BM392983 50071-2-4 BM394450 50072-2-4 AQ588780 CITBI-E1-AZ539947 ENTDL61TF AZ539947 ENTDE20TF BX386038 BX386038 AG046758 Pan trogl	Drosophi CH261-74 L536976	Drosophi 223PbH05 Homo sap Homo sap 11a17c06	ENTCZ15 BX46131 WHE1257 BX39812 nbxb004 CH261-1	7028c02. 7142f07. 7142f07. 01-M-F0 063PbH0 PUGC069	BX43646 Tetraod CH261-9

ALIGNMENTS

MEDLINE PUBMED COMMENT	TITLE		REFERENCE AUTHORS	VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 BF373406 LOCUS DEFINITION ACCESSION
Proc. Natl. Acad. Sci. U.S.A. 9/ (/), 3491-3496 (2000) 20202663 10737800 Contact: Simpson A.J.G. Contact: Simpson A.J.G. Laboratory of Cancer Genetics Ludwig Institute for Cancer Research Ludwig Institute for Cancer Research Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,	Simpson, A.J. Shotgun sequencing of the human transcriptome with ORF expressed sequence tags	Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOllveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and	<pre>Bukaryote; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 167) Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,</pre>	BF373406.1 GI:11335431 EST. Homo sapiens (human) Homo sapiens	BF373406 IL2-FT0159-070800-120-H01 FT0159 Homo sapiens cDNA, mRNA sequence. BF373406

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REFERENCE
AUTHORS
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AA578773
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence.
AA578773
                                                                                                                                                                                                                                                                                                                         Mammalia; Butheria; Primates; Catarrnin; Hom 1 (bases 1 to 402)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA578773
402 bp mRNA linear EST 12-SEP-199
nh24a04.s1 NCI_CGAP_Pr1 Homo sapiens cDNA clone IMAGE:953262, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: puc 18 forward High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtm12.pl7t1=IL2&t2=IL2-FT0159-
070800-120-H01&t3=2000-08-07&t4=1)
CDNA Library Preparation: David B. Krizman, Ph.D. CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, DNA Sequencing by: Washington University Genome Sequencing Clone distribution: NCI-CGAP clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 565 Std Error: 0.00
                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Fax: +55-11-2707001
                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                National Cancer Institute, Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                        Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=Torgan: prostate_tumor; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-llbrary was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone
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/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ity sequence stop: 167.
Location/Qualifiers
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Pred. No. 2.3e-35;
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                                                                          Sequencing (information
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RESULT 3
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133; Conserv
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1 (bases 1 to 489)

1 (bases 1 to 489)

McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, J., Wylie, T., McCarter, M., Marra, M., Hillier, L., Kucaba, T., Theising, B., Bowers, Y., Dante, M., Mitter, E., Bennett, J., Franklin, C., Teagareishvili, R., Gibbons, M., Kennedy, S., Maguire, L., Beck, C., Underwood, K., Steptoe, M., Allen, M., Person, B., Swaller, T., Harvey, N., Schurk, R., Kohn, S., M., Allen, M., Person, B., Cardenas, M., McCann, R., Waterston, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BG224987
BG224987.1
EST.
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Location/Qualifiers
                                                           The Washington Univ. Nematode EST Project, Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Lo
                                                                                                                                                                                                                                                                                                                                                                                                              Strongyloides stercoralis
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BG224987 489 bp mRNA linear EST 09-MP kp52h05.y1 TBN95TM-SSFH Strongyloides stercoralis cDNA 5' sin to WP:C08B11.7 CE01477 HYPOTHETICAL 37.7 KD PROTEIN C08B11.7
                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                   Unpublished Contact: McCarter JP
                                                                                                                                                                                The Washington Univ. Nematode EST Project, 1999
                                                                                                                                                                                                                                                                                                                                                                                           Panagrolaimoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected, histologically normal prostate epithelial cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
est@watson.wustl.edu
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/note="Vector: pAMP10; Si
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/lab_host="DH10B"
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|mol_type="mRNA"
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Pred. No. 1.9e-22;
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library was constructed

λÃ Dr.

Thomas Nutman and colleagues

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REFERENCE
AUTHORS
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BX442677/c
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Best Local Similarity
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                                                                                                                                                                  http://www.genoscope.cns.fr/
cgi-bin/cluster.cgi?seq=CSODF035AH01NP1&cluster=1534.f. Contact
Feng Liang Email: fliang@lifetech.com URL:
Feng Liang Email: fliang@lifetech.com URL:
Feng Liang Email: fliang@lifetech.com URL:
                                                                                                                                                                                                                                                                       Email: segref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen. This sequence belongs to sequence cluster 1534.f
more information about this cluster, see
                                                                                                                                                                                                                                                                                                                                                                         Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 1016)
Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
Full-length cDNA libraries and normalization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BX442677 Homo sapiens FETAL BRAIN Homo CSODF035YCO1 3-PRIME, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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Location/Qualifiers
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/note="Vector: Lambda Uni-ZAP XR (Stratagene); Site 1:
/note="Vector: Lambda Uni-ZAP XR (Stratagene); Site 1:
/note="Vector: Lambda Uni-ZAP XR (Stratagene); Site 1:
/note="Vector: Lambda Uni-ZAP XR (Stratagene); Site 2: XNoI; mRNA was purified from 4 x 1085
/filariform larvae which had been isolated from infected
/numans. cDNA was constructed and, using adaptors, was
/cloned unidirectionally into the vector from the EcoRI
/site to the XNoI site. The library has an unamplified
/titer of 1.5 x 1086 pfu/ml and an amplified, undiluted
/titer of 7 x 1089 pfu/ml. The average insert size of the
/numanplifed library is 975 bp (range, 500-1500)."
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/mol_type="mRNA"
/strain="Filariform larvae obtained from humans"
/clone="CS0DF035Y001"
                                              organism="Homo sapiens"
/mol_type="mRNA"
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/lab_host="XL-1 Blue MRF′ (Stratagene)"
                            db_xref="taxon:9606"
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602 KKKAGKAAKAAKAAADAAADAAKAKAAGKKKAAAAARKDDKKKKNAAAAAAAAAAA
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EX436460 Homo sapiens THYMUS Homo sapiens cDNA clone CSOCAP001YL10
5-PRIME, mRNA sequence.
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Location/Qualifiers
                                                                                       40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1098)
Li W.B., Gruber,C., Jessee,J. and Polayes,D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bmail: seqref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen.Contact: Peng Liang Email: filang@lifetech.com
http://fulllength.invitrogen.com/ Invitrogen Corporation 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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BP 191 91006 EVRY cedex - France
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                                        GTGAGGGAAGGACATTAGAAAATGAATTGATGTGTTCCTTAAAGGATGGGCAGGAAAACA
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                                                                                       Conservative
                                                                                                                                                                                      /tissue_type="THYMUS"
/clone_Tib="Homo sapiens THYMUS"
/clone_Tib="Homo sapiens THYMUS"
/note="Vector: pcMVSPORT 6; lst strand cDNA was primed with a NotI-oligo(dT) prImer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned i the Not I and EcoRV sites of the pcMVSPORT 6 vector.
Library was not normalized."
289 c 225 g 229 t 299 others
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/note="Organ: brain; Vector: pCMVSPORT_6; 1st strand cDNA
was primed with a NotI-oligo(dT) primer. Five prime end
enriched, double-strand cDNA was digested with Not I and
cloned into the Not I and ECORV sites of the pCMVSPORT 6
vector. Library was not normalized."
210 c 197 g 233 t 212 others
                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                       clone="CS0CAP001YL10"
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22.7%;
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                                                                        103 GAAGGACATTAGAAAATGAATTGATGTGTTCCTTAAAGGATGGGCAGGAAAACAGATCCT
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137L10 of
                                                                                                                                                                                                                                                                                                                                                                                     Submitted (12-ARR-2000) Genoscope - Centre National de Sequencage : BP 191 1006 EVKY Cedex - FRANCE (E-mail : seqref@genoscope.cns.fr - Web : www.genoscope.cns.fr)

This sequence is a single read and was generated as part of a large scale clone-end sequencing project of the Tetraodon nigroviridis genome. For more information, please take a look at http://www.genoscope.cns.fr/Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roest Crollius, H., Jaillon, O., Dasilva, C., Ozouf-Costaz, C., Fizames, C., Fischer, C., Bouneau, L., Billault, A., Quetier, F., Saurin, W., Bernot, A. and Weissenbach, J. Characterization and repeat analysis of the compact genome of reshwater pufferfish Tetraodon nigroviridis Genome Res. 10 (7), 939-949 (2000)
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Nat. Genet. 25 (2), 235-238 (2000)
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Tetraodon nigroviridis
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GTTGTGGATATTTATTTGAACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCAT
                                      DAAASAKAWTAKACAAKGTWKTGTATTATAKACAAAAWASGTGTATRTAWWAWTATTGT
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/note="Genoscope sequence
66 c 233 g 211
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/mol_type="genomic_DNA"
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library G from Tetraodon
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                                                                                                                 14.4%; Score 40; DB;
41.3%; Pred. No. 2.6;
tive 27; Mismatches
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nigroviridis
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1 of clone
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BASE COUNT
ORIGIN
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CC241440/c
                                          RESULT 8
BI541137
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ORGANISM
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                                                                                                     816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        223
                                                                                                                                 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           959 AAAATAKTATAAAAAAATATSABWTTAAAT
BI541137
454735 MARC :
BI541137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High quality sequence start: 48
High quality sequence stop: 704.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Richard K. Wilson
Genome Sequencing Center
Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 1051)
Kremitzki,C., Higginbotham,J.,
Warren,W., Graves,T., Mardis,E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC241440.1 GI:30568103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Archosauria; Aves; Neognathae; Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gallus gallus (chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CH261-96M14_Sp6.1 CH261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: Sp6 ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: submissions@watson.wustl.edu
Insert Length: 182000 Std Error: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gallus gallus BAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                                          TACCAATGAGAGGAAAACAGACGAGAAAAT
                                                                                                                                                              TAAAAGGTGATTTGGAGTTGAAAAAACAGAAATCCAAGGGAATATTTCAATAGGGGGGGAA
                                                                                                                                                                                            TGAACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAA
                                                                                                                                                                                                                         TTÄÄTTGGTTTTTTCCTAÄGGTAÄAAGGTTTATÄÄTGGTACCCCCCTGCCATTTTTGGTT
                                                                                                     GCTTATGAAAAAAATTCAAAGGTTCA
                                                                                                                                 ACAGACGAGAAAATCTTGATGGCTTCA
                                                                                                                                                                                                                                                                                                                                                         351
                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                       CH261 Female Chicken library - for library and clo ordering information: http://www.chori.org/bacpac" a 205 c 203 g 292 t
                                                                                                                                                                                                                                                                                                                                                                                    /note=\overline{}^{\mathrm{T}}Vector: pTARBAC2.1; Site 1: EcoRI; Site 2: EcoRI; CH261 Female Chicken library - \overline{}for library and clone
                                                                                                                                                                                                                                                                                                                                                                                                                                            /strain="Red Jungle Fowl"
/db_xref="taxon:9031"
/clone="CH261-96M14"
/sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Gallus gallus"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                 /clone
                                                                                                                                                                                                                                                                                                                                                                                                                               cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .1051
                1BOV
                                                                                                                                                                                                                                                                                               13.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                              line="UCD001, inbred 256"
                Bos
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                taurus
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                                                                                                                                                                                                                                                                                               Score 38.2;
Pred. No. 8
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                393 bp mRNA
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                                                                                                                                  . 265
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. and Wilson,
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genomic clone
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on,R.
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                               linear
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CH261-96M14,
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                               EST 30-AUG-2001
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                                                                                                                                                                                                                                                                                 Gaps
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REFERENCE
AUTHORS
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AA815526
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   AUTHORS
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                                                                                                                                                                                                                                                                                             277
                                                                                                                                                                                                                                                                                                                                142 ATGGGCAGGAAAACAGATCCTGTTGTGGATATTTATTTGAACGGGATTACAG 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 393)
Smith,T.P.L., Grosse,W.M., Freking,B.A., Roberts,A.J., Stone,R.T., Casas,E., Wray,J.E., White,J., Cho,J., Fahrenkrug,S.C., Bennett,G.L., Heaton,M.P., Laegreid,W.W., Rohrer,G.A., Chitko-McKown,C.G., Pertea,G., Holt,I., Karamycheva,S., Liang,F., Quackenbush,J. and Keele,J.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                            65,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea Bovidae; Bovinae; Bos.
                                                                                                                                                                vp28c02.r1 Barstead
clone IMAGE:1077986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plate: 119 row: I column: 22
Seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BACKWARD: GTTTTCCCAGTCACGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single pass sequencing. Bases (v0.980904.e. Vector identified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: smith@email.marc.usda.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 402 762 4366
Fax: 402 762 4390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
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EST.
   Marra,M.,
                                                                      Mus musculus
                                                                                                                             AA815526.1
                                                                                                                                                                                                  AA815526
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome Res. 11 (4), 626-630 (2001)
                                    Mammalia;
                                                     Eukaryota; Metazoa;
                                                                                                                                              AA815526
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                                                                                       Mus musculus (house
                                                                                                                                                                                                                                                                                             CTATACTGGGAAACAGCTGGTACTGGGAAAACTTTTCAGAAATGGATAACAG
                                                                                                                                                                                                                                                                                                                                                                                                    TGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATGAATTGATGTGTTCCTTAAAGG 141
                                                                                                                                                                                                                                                                                                                                                                    TGCCCTANAGAGTTTGAAGGTGAAATACATTGGGAAACAAATGATGAACAAATGAAAGG
                   (bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRimers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      112
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pCMV SPORT6; Site 1: NotI; Site 2: SalI; Library made from pooled tissue from lymph node, ovary, fat, hypothalamus, and pituitary."
77 c 116 g 87 t 1 others
                 Eutheria;
1 to 206)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="pooled"
/lab_host="DH10B"
/clone_lib="MARC_1BOV"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:9913"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Bos taurus"
(mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
Hillier, L., Allen, M., Bowles, M.,
                                                                                                                             GI:2885122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GI:15382249
                                                                                                                                                                                                                                                                                                                                                                                                                                                          13.5%;
                                Chordata;
Rodentia;
                                                                                                                                                            206 bp mRN mouse proximal col 5', mRNA sequence.
                                                                                         mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 37.4;
Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                  Sciurognathi;
                                                     Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         called and alt_
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                                                                                                                                                                                 mRNA linear
colon MPLRB6 Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                            47;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 393;
Dietrich, N., Dubuque, T.,
                                    Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      trimmed with phred a with the -minscore 18
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                                                                                                                                                                                   musculus
                                                                                                                                                                                                EST 13-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 10
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Best Local
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                                                                        AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                            247
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                                                                                                                                                                                                                                                                                                                                                         186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   66
1 (bases 1 to 384)
Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T.,
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
Waterston,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fax: 314 200 10.2
Email: mouseest@waitson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylle,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                           vt42f07.r1 Barstead mouse proximal colon MPLRB6
clone IMAGE:1165765 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished Contact: Marra M/Mouse EST Project
                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                          Mus musculus
                                                                                                                                                            Mus
                                                                                                                                                                             AA710451.1
EST.
                                                                                                                                                                                                                                                                    AA710451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The WashU-HHMI Mouse EST Project
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                                                                                                                                                                                                                                                                                                                                                                                          GAAAA 251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer: -28ml3 rev2 ET from Amersham.
                                                                                                                                                           musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone lib="Barstead mouse proximal colon MPLRB6"
/note="Vector: pT/T3D-Pac (Pharmacia) with a modified
polylinker; Site I. EcoRI, Site 2: NotI, lst strand cf
was primed with a Not I - oligo(dT) primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="IMAGE:1077986"
/dev_stage="7 day juvenile"
/lab_host="DH10B"
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strain="FVB/N"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            187 ATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAACAGACGA 246
                                                                                                                                                                                                                                              397 bp mRNA linear UI.M-FOO-cdo-m-15-0-UI.rl NIH_BMAP_FOO Mus musculus IMAGE: 6831256 5', mRNA sequence.
CA752332
CA752332.1 GI:25582740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: -28m13 rev2 ET from Amersham High quality sequence stop: 363.
Location/Qualifiers
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Jim Lin, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
                                                                                                     Mammalia; Eutheria; Rodentia; Sciurognathi; Mu
1 (bases 1 to 397)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
                                                                                                                                                                                                   Mus musculus
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4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Marra M/Mouse EST Project WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MGI:631677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                          Jnpublished
                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                   Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: mouseest@watson.wustl.edu
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314 286 1810
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67 c 65 g 88 t
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/lab_host="DH10B"
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'clone="IMAGE:1165765"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .384
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56.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double-stranded cDNA was ligated to Eco RI adaptors CGGATCCTTG), digested with Not I and cloned into the
                                                                                                                                                             Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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Pred. No. 14;
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RESULT 12
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                                                  Plasmodium berghei

Bukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.

Bukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.

1 (bases 1 to 811)

Carlton, J.M.-R. and Dame, J.B.

The Plasmodium vivax and P. berghei gene sequence tag projective parasitol. Today (Regul. Ed.) 16 (10), 409 (2000)
                                                                                                                                                                                                                                                           berghei cDNA 3', mRNA
BF299229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
This clone was contributed by the Brain Molecular Anatomy Project
Parasitol. Today (Regul. Ed.) 16 (10), 409 (2000)
Contact: Dame JB
Department of Pathobiology, College of Veterinary Medicine
University of Florida
                                                                                                                                                                                                  Plasmodium berghei
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                                                                                                                                                                                                                                             BF299229.1 GI:13952844
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence: 217-329, >POLY_A#Simple_repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The following repetitive elements were found in this cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAACAGACGAGAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACTCTTGGCCAAAATTTAGAAGATGCTGTGAATATCATTTTGAACTTGTGTAAATATATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="embryo 12.5dpc"

//dev_stage="embryo 12.5dpc"

//lab_host="DH10B (T1 phage resistant)"

//clone_lib="NIH_BMAP_FOO"

//clone_lib="NIH_BMAP_FOO"

//note="Organ: Brain; Vector: pyx- Asc; Site_1: EcoR I;

Site_2: Not I; The library was constructed according

Bonaldo, Lennon and Soares, Genome Research, 6:791-806,

1996. Denatured RNA was size fractionated on a 1% agazose

gel. First strand cDNA synthesis was primed with oligo-dT

primer containing a Not I site. Double strand cDNA was

size selected according to mRNA size fraction, ligated

with EcoR I adaptor, digested with NotI and then cloned

directionally into pYX-Asc vector. The library tag

sequence located between the NotI site and the polyA tail

is TGAAGAGACC. This library was created for the University

Iowa Brain Anatomy Project (BMAP): 'Gene Discovery in the

Developing Mouse Nervous System', supported by National

Institute of Mental Health (NIMH), Hemin Chin, Ph.D.,

program coordinator."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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/tissue_type="whole brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mol_type="mRNA"
/strain="C57BL/6"
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Pred. No.
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rles Yowell and Jane
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. 14;
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                                                                      ne sequence tag projects
409 (2000)
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on Plasmodium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   69 GAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATGAATTGATG
                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

1 (Dases 1 to 804)

Whitelaw, C.A., Quackenbush, J., Van Aken, S., Utterback, T., Resnick, A., Fraser, C.M., Yuan, Y., San Miguel, P., Ma, J. and Bennetzen, J.
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Tel: 352 392 4700
Fax: 352 392 9704
                            TIGR
9712 Medical Center Drive, Rockville,
                                                                                                          Unpublished
Other_GSSs: PUGCO69TD
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PUGCO69TB ZM_0.6_1.0_KB Zea mays
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                                                                                  Contact: Cathy Whitelaw
                                                                                                                                                     Maize Genomics Consortium
                                                                                                                                                                                                                                                                                                                      Zea mays
                                                                                                                                                                                                                                                                                                                                                                                        BZ819014.1 GI:29033836
                                                                                                                                                                                                                                                                                                                                                                                                                                genomic survey sequence
                                                                                                                                                                                                                                                                                                                                           Zea mays
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301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone lib="Pb cDNA #20, Charles Yowell and Jane Carlton"
/note="Vector: pBluescript II vector DNA, excised from
Lamda ZAP II; Site 1: ECORI, Site 2: XhoI; Total RNA was
extracted from asynchronous blood stage forms of the
cloned ANKA isolate of P. bezghei grown in laboratory
Swiss white mice. Contaminating host white cells had
previously been removed using a novel blomagnetic bead
protocol (J. Carlton et al., manuscript in preparation).
Polya+ RNA was extracted and reverse transcribed using an
oligo dT-XhoI primer. Second strand cDNA was prepared
using RNase H and DNA polymerase I. ECOR I adaptors were
ligated to the cDNA, and it was digested with XhoI.
Fragments were size selected, and those between 1-5 kb
ligated into ECORI /XhoI digested vector."
                  301-838-5843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   damej@mail.vetmed.ufl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="asynchronous blood stage"
/lab_host="Swiss white mice"
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/dev_stage="asynchron
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Pred. No. 17;
0; Mismatches 65
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AUTHORS
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 219 GCATTACCAATGAGAGGAAAACAGACGAGAAAATCTTGATGGCTTCAC 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       159 TCCTGTTGTGGATATTTATTTGAACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGA 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99 GAGGGAAGGACATTAGAAAATGAATTGATGTGTTCCTTAAAGGATGGGCAGGAAAACAGA 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   86;
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Class: sheared ends
                                                                                                                                                                                                                 Seq primer: TF
Class: sheared ends.
                                                                                                                                                                                                                                                                                                                                                             Unpublished
Other_GSSs: PUGCO69TB
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                                                                                                                                                                                                                                                                                    9712 Medical Center Drive,
Tel: 301-838-5843
                                                                                                                                                                                                                                                                                                                                                                                                                Whitelaw, C.A., Quackenbush, J., A., Fraser, C.M., Yuan, Y., San
                                                                                                                                                                                                                                                   Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                     Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                            TIGR
                                                                                                                                                                                                                                                                                                                                         Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                                                                                                                 Maize Genomics Consortium
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             /db_xref="taxon:4577"
/clone="ZMMBTa337L17"
/clone="EMMBTa337L17"
/clone="Vector: pCR4-TOPO; Site_1:
/note="Vector: pCR4-TOPO; Site_1:
COT selected genomic DNA library"
a 157 c 158 g 268 t
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                                                                                                                     /mol_type="genomic DNA"
/strain="B73"
                                                                                                                                          /organism="Zea mays"
/mol_type="genomic D
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/clone lib="ZM 0.6 1.0 KB"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; 0.6-1.0 kb high
COT selected genomic DNA library"
141 c 144 g 273 t
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|mol_type="genomic D
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Pred. No. 19;
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                                                                                                                                                                                                                                                                                                         Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                J., Van Aken, S., Utterback, T., Resnic San Miguel, P., Ma, J. and Bennetzen, J
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Query Match

13.2%;

Score 36.8;

DB 29;

Length 880;

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168 GGATATTTATTTGAACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCA 227

Search completed: February 1, 2004, 10:40:53 Job time : 1319 secs

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seq length: 0 seq length: 2000000000
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ABL95270	AAF86949	AAH02871	AAH85120	AAH93806	AAS63899	AAA06690	AAF30667	ID	SUMMARIES
Human P710P cDNA s	Human P710P invent	Prostate tumour an	Human prostate-spe	Human prostate-spe	Human prostate cDN	Human immunogenic	Human differential	Description	

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AAA06545	0	ACA59834	ABL95397	ABK92196	AAH93861	AAS64026	AAV62430	ശ	526	AAF86946	AAH02868	AAH85117	AAH93803	AAS63896	AAA06687	ACA59706	ABL95269	AAF86948	. AAH02870	AAH85119	AAH93805	AAS63898	AAA06689	ACA59705	ABL95268	694	AAH02869	AAH85118	AAH93804	AAS63897	AAA06688	AAV62427	AAV62428	881	AAV62429	ACA59707
Human immunogenic	fferenti		Human DD3 cDNA seq	Prostate cancer-as	Gene DD3 cDNA sequ	Human cDNA encodin	Prostate cancer an	Prostate cancer th		Human P710P invent	Prostate tumour an	Human prostate-spe	Human prostate-spe	Human prostate cDN	Human immunogenic	Ġ)	CDNA	Human P710P invent	ate tumour	prostate-s			Human immunogenic	Prostate cancer th	P710P	P71	Prostate tumour an	prostate-s		Human prostate cDN	Human immunogenic	Prostate cancer an	Prostate cancer an	7.	Prostate cancer an	Prostate cancer th

AAF30667 standard; cDNA; 278 BP. 11-JUN-2001 (first entry)

ALIGNMENTS

Human differentially expressed PCA3 cDNA (short form).

human; PCA3; prostate cancer; antigen; benign prostatic hyperplasmia; differential expression; diagnosis; gene therapy; chromosome 9; .83

Homo sapiens

29-SEP-1999; 29-SEP-2000; 2000WO-CA01154. 05-APR-2001. WO200123550-A2. exon 99US-0156594. /*tag= a /number= "3" 27..278 /*tag= b /number= "4a" Location/Qualifiers

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RESULT 2
AAA06690/c
ID AAA066
XX AAA066
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XX Human
XX Human;
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Matches 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of an RT-PCR-amplified fragment of human prostate cancer antigen 3 (PCA3) mRNA. A 2nd amplified PCA3 sequence (see AAF30666) includes an additional 27 bp, inserted between exons 3 and 4a. The shorter form RNA is associated with prostate cancer whereas the longer form RNA is associated with a non-malignant prostatic state, such as benign prostatic hyperplasia. Based on the differential expression of these 2 PCA3 RNA species, protocols for the diagnosis of prostate disease are provided, including a method of diagnosing the presence or predisposition to develop prostate cancer in a patient. Also provided are therapeutic methods that use a nucleic acid encoding a differentially expressed PCA3 mRNA molecule, an antisense sequence, a protein encoded by a differentially expressed pCA3 mRNA molecule, an antisense sequence, a protein encoded by a differentially expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel nucleic acid encoding differentially expressed prostate cancer antigen 3 mRNA containing additional sequence giving rise to long PCA3 mRNA, useful for diagnosis of mammal afflicted with prostate cancer -
                                                                 WO200004149-A2
                                                                                                                                                                                                                         Human; prostate cancer; diagnosis; tumour; gene therapy; detection;
                                                                                                                                                                                                                                                                                    Human immunogenic prostate tumour protein cDNA sequence SEQ ID NO:471
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                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                  (first entry)
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27-JAN-2000.

RESULT 3
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ID AAS638
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AC AAS638
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AC AAS638
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DE Human 1

AAS63899

standard; cDNA; 812

383 180 443 120 503 60

Human prostate cDNA sequence #433

29-JAN-2002 (first entry)

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23-SEP-1998;
15-JAN-1999;
15-JAN-1999;
09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             immunogenic portion of a prostate tumour protein (PTP). The polypeptides and polynucleotides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polypeptides and polynucleotides encoding them, antigen presenting cells which express the polypeptides, antibodies against the polypeptides and vaccines comprising them can be used for inhibiting the development of prostate cancer in a patient. The polypeptides can be used to generate antibodies or anti-idiotypic antibodies for passive immuno therapy. A portion of the polypucleotides encoding the polypeptides can be used as a probe or composite the expression of the polypeptides. AAA06241 to AAA06691 and AAV82000 to AAV82020 represent sequences used in the exemplification of the polypeptides.
                                                                                                                                                                                                                                                                                                                                                                      Sequence 812 BP; 198 A; 189 C; 169 G; 256 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 262; 263pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide useful for treating comprises an immunogenic portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-171268/15
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14-JUL-1998;
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                    AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
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AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
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                                                                                                                       Harlocker
                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention describes isolated polypeptides, operation of a prostate tumour protein (PTP).
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99US-0232149.
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99US-0288946.
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Pred. No. 4.7e-76;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and diagnosing prostate cancer prostate tumor protein -
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286
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09-MAY-2000; 2000US-0568100.
12-MAY-2000; 2000US-0593793.
13-UUN-2000; 2000US-0693793.
27-JUN-2000; 2000US-0636215.
29-AUG-2000; 2000US-0657279.
06-SEP-2000; 2000US-0657279.
02-OCT-2000; 2000US-0685166.
                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                  response, ....detecting cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Хu J,
                                                                                                                                                                                                                                                                                                                                                                          Sequence 812 BP; 198 A; 189 C; 169 G; 256 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 386-387; 579pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to isolated prostate-specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human prostate-specific polypeptides and polynucleotides useful diagnosis and treatment of cancer, especially prostate cancer.
 323
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                                                                                                                                                       TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCCGCTTGTGAGGGAAAGGACATTAGAAAATG
                                                                                                                                                                                                                                                                   CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCCGGCCGCCCATCTTGGG
                     AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                            AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                         AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGAAAAC 240
                                                                                                                        TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                                                                                                              CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       on DC, Mitcham JL, Harlocke
Retter MW, Stolk JA, Day C
g A, Skeiky YAW, Hepler WT,
                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer; ss; cytostatic; immunostimulant; tumour.
                                                                                                                                                                                                                                                                                                                            95.7%;
99.6%;
                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                            Score 266; DB 22;
Pred. No. 4.7e-76;
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/A, Day CH, Vedvick
fepler WT, Henderson
                                                                                                                                                                                                                                                                                                               0;
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dvick TS,
lerson RA;
                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                            812;
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Carter D;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                The present invention describes polynucleotide sequences (I) which encode prostate-specific proteins (II). (II) and (II) have cytostatic activity, and can be used in vaccine production and gene therapy. (I), (II), antibodies to (II), fusion proteins comprising (II), and isolated T cells prepared using (I) or (II) are used treat cancer in a patient. The cancer that is diagnosed or treated is particularly prostate cancer. (I) and (II) can be used in the detection of cancer in a patient. (I) and (II) can be used for monitoring the progression of cancer in a patient. (II) can be used for monitoring the progression of cancer in a patient. (II) and (II) can also be used to improve diagnostic and therapeutic methods for prostate cancer. They can indicate the level of metastasis as well as the prostate volume. AAM93357 to AAM9344 and AAM0115 to AAM01318 represent polynucleotide and amino acid sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                        Sequence 812 BP; 198 A; 189 C; 169 G; 256 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polynucleotide encoding a prostate-specific protein, diagnosing, monitoring and treating prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; prostate cancer; prostate-specific; diagnosis; vaccine; cytostatic; gene therapy; metastasis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 385; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kalos MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-JAN-2000; 2000US-0483672.
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                                                          121
 181
                                                                                                                                                                                                              277;
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                                                                                                                      13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dillon DC, Mi
MD, Fanger GR,
                                                                                                                                                                      1 CAGGAAGCACAAAAAGGAAGCACAGAGATCCCCTGGGAGAAATGCCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Meagher MJ;
                                                                                                         TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGAAGGACATTAGAAAAATG
                                                TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                    CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
 AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                             Conservative
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GR, Day CH,
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99.6%;
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                                                                                                                                                                                                                            Score 266; DB 22;
Pred. No. 4.7e-76;
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                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                        0 other;
                                                                                                                                                                                                              0
                                                                                                                                                                                                                                          Length 812;
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AAH85120/c
ID AAH85120 standard; cDNA; 812
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                                                                                                                 Query Match
Best Local S
Matches 277
                                                                                                                                                                                                 The present invention describes an isolated polypeptide (P1) comprising at least an immunogenic portion of a prostate-specific protein, or its variant. Also described are polynucleotides (N1) encoding (P1). (P1) and (N1) have cytostatic activity and can be used in vaccine production. The polypeptides, nucleic acids and antibodies from the present invention are useful in the diagnosis and therapy of prostate cancer. Prostate specific genes P704P, P712P, P774P, P775P and B305D are located in a genomic region on chromosome 22q11.2 known as the Cat Eye Syndrome region. Prostate specific antigen (PSA) P501S was located on chromosome 1. AAH84671 to AAH85143 and AAG99000 to AAG9907 represent polynucleotide and polypeptide sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; prostate cancer; therapy; diagnosis; cat eye syndrome; chromosome 22q11.2; prostate-specific protein; chromosome 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated polypeptide comprising at least prostate-specific protein, useful in the prostate cancer -
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                                                                                                                                                                          Sequence 812 BP; 198 A; 189 C; 169 G; 256
                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 284-285; 325pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-308785/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kalos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-NOV-1999;
18-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prostate specific antigen; PSA; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH85120;
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                                                                                                                 Local Similarity
les 277; Conserv
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                                                          562
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MD, Retter MW, Stolk JA,
                     TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGGAAGGACATTAGAAAATG 120
                                                        CAGGAAGCACAAAAGGAAGCACAGAGATCCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
                                                                                      CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
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                                                                                                                   Conservative
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99US-0443686
                                                                                                                              95.7%;
99.6%;
                                                                                                                   0; Mismatches
                                                                                                                              Score 266;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harlocker SL, Jiang Y,
A, Day CH, Skeiky YAW,
                                                                                                                               DB 22;
1.7e-76;
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                                                                                                                                             Length 812;
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Matches Query Match

Local Similarity nes 277; Conserv

Conservative

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Mismatches

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Gaps

95.7**%**; 99.6**%**;

Score 266; DB 22; Pred. No. 4.7e-76;

Length 812; Indels

1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCCGGCCGCCATCTTGGG

CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG

TCATCGATGAGCCTCGCCCTGTGCCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG

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                        The present invention describes an isolated polypeptide (I) comprising at least an immunogenic portion of a prostate tumour antigen protein or its variant. (I) have cytostatic activity and can be used in vaccine production. (I) prostate tumour antigen polynucleotides, an antigen presenting cell (APC e.g. a dendritic cell) that expresses (I), and a pharmaceutical composition containing (I) are useful for inhibiting the development of cancer in a patient. Antibodies specific for prostate specific proteins and oligonucleotides that hybridise to a polynucleotide that encodes a prostate specific protein are useful for detecting the presence or absence of a cancer or monitoring the progression the progression of a cancer, especially prostate cancer. AAH02422 to AAH2872, AAB74798 to AAB74821 and AAB74830 are sequences used in the exemplification of the present invention.
 Sequence 812 BP; 198 A; 189 C; 169 G; 256 T; 0 other;
                                                                                                                                                                                                                                                                     Prostate specific protein and its encoding polynucleotide, useful for the treatment and diagnosis of prostate cancer -
                                                                                                                                                                                                                                                                                                                    WPI; 2001-245062/25
                                                                                                                                                                                                                                                                                                                                                                                                         04-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                       04-OCT-2000; 2000WO-US27464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; prostate tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prostate tumour antigen cDNA sequence for P710P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-APR-2001
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                                                                                                                                                                                                                                          4; Page 267; 276pp; English.
                                                                                                                                                                                                                                                                                                                                                Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
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                                                                                                                                                                                                                                                                                                                                                                                                           99US-0157455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA;
                                                                                                                                                                                                                                                                                                                                                 Reed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           n; prostate tumour; therapy; diagnosis;
cytostatic; vaccine; ss.
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RESULT 7
AAF86949/c
ID AAF869
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XX Human
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XX Chromo
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XX 12-APR
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XX PO170-
PF 04-OCT
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Best Local
                                                                                                                                                     Matches 277;
                                                                                                                                                                                                                                                                    The present invention describes compositions comprising peptides derived from the Wilm's tumour protein WT1 and methods for their use in treating malignant diseases. Peptides derived from both the murine and human WT1 proteins are provided. The human WT1 gene is found on chromosome 11p13, and the protein was shown to be a zinc finger transcription factor. The immunogenic peptides of the invention are particularly useful in the diagnosis and treatment of cancer and leukaemia. The present sequence is a coding sequence used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide comprising part of the Wilm's Tumour gene product sequence is used in the diagnosis and treatment of malignant diseases e.g. leukemia and cancer associated with WTI \, -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; mouse; immunotherapy; cancer; leukaemia; WT1; Wilm's tumour gene; chromosome 11p13; zinc finger transcription factor; ss.
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                                                                                                                                                                                                                               Sequence 812 BP; 198 A; 189 C; 169 G; 256 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 220; 228pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-328324/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-OCT-1999;
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                                                                                                                                                                        Local Similarity
502
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                                  61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORIXA CORP
                                                                                          CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                  TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG 120
                                                                           CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                   95.7%;
ilarity 99.6%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0157459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cheever MA,
                                                                                                                                                   Score 266; DB 22;
Pred. No. 4.7e-76;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    278
                                                                                                                                                                                        Length 812;
                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         345.
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                                                                                                                                                     Gaps
                                                                                                              60
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                                                                         503
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(WANG/)

CARTER D.
LI S X.
WANG A.
SKEIKY Y A W.
HEPLER W T.

LISX/)

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RESULT 8
ABL952
XX ABL952
XX ABL952
XX ABL952
XX ABL952
XX ABL952
XX ABL952
XX ABL952
XX Human
XX Human;
XX Human;
XX US2002
XX J-FEB
PN US2002
XX 12-FEB
PR 25-FEB
PR 25-FEB
PR 25-FEB
PR 12-JAN
PR 13-JUL
PR 13-JUL
PR 13-JUL
PR 13-JUL
PR 13-JUL
PR 13-JUN
PR 11-AUG
PR (AUGJ/
PR (AUGJ/
PR (AUGJ/
PR (CART//
PR (
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27-MAR-2000;
09-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                         13-JUN-2000;
27-JUN-2000;
10-AUG-2000;
29-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-199;
12-NOV-1999;
18-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JAN-1999;
09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-FEB-1998;
25-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-FEB-1997;
01-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cancer; prostate cancer; vaccine; cytostatic; immunostimulant; gene therapy; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human P710P cDNA sequence SEQ ID NO 471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL95270;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL95270 standard; cDNA;
                                                                             (HARL/)
(JIAN/)
(KALO/)
(FANG/)
(RETT/)
(STOL/)
(DAYC/)
(CART/)
                                                                                                                                                                                                                                                              (XUJJ/)
                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2000;
02-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-JUL-1998;
23-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2002022248-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                10-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
                                                                                                JIANG Y.

KALOS M D.

KALOS M D.

RETTER M W.

RETTER M W.

STOLK J A.

DAY C H.

VEDVICK T S.
                                                                                                                                                                                                                                                                MITCHAM J
                                                                                                                                                                                                                                             MITCHAM J L.
HARLOCKER S L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                               2000US-0593793.
2000US-0605783.
2000US-0636215.
2000US-0651236.
2000US-0657279.
2000US-0679426.
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2000US-0536857.
2000US-0568100.
2000US-0570737.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0030607.
98US-0115453.
98US-0159812.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0443686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0352616
99US-0439313
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0232149
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RESULT 9
ACA59707/c
ID ACA597
XX ACA597
XX ACA597
XX Prosta
XX Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
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Best Local S
Matches 277
                                    17-APR-2000;
04-OCT-2000;
28-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                      immunogen; cancer; vaccine; gene therapy; cytostatic; prostatic antigen; PSA; prostatic acid phosphatase; PAP; prostate acceptable acid phosphatase; PAP; prostate acceptable ac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The
thei
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Fanger
Li SX,
                                                                                                                                                                 29-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           progression
                                                                                                                                                                                                                                                                                                 US2002192763-AJ
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prostate cancer therapy associated cDNA #450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACA59707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 812 BP; 198 A; 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 471; 87pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New prostate-specific polynucleotides diseases, in particular prostate cance progression of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HEND/) HENDERSON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancers, pa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         323
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GR, Retter MW, Stolk JA,
Wang A, Skeiky YAW, Hep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention provides prostate-specific coding sequences and ed proteins. These can be used in the diagnosis and treatme particularly prostate cancer. The present sequence is a cf n the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                    2000US-157455P.
2000US-0679272.
2001US-0822827.
                                                                                                                                                                    2001US-0895793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ×
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95.7%;
99.6%;
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Pred. No. 4.7e-76;
0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  278
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fusion protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       812;
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Carter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment is a cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     324
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                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                         The invention describes a fusion protein comprising at least one amino acid sequence of immunogenic portions of any of the 3 sequences not defined in the specification, or sequences having at least 70 or 90 % sequence identity to any one of the 35 sequences defined in the USPTO web site, which is encoded by any of the 4 nucleotide sequences not defined in the specification. The fusion protein, composition and methods are useful for diagnosing, preventing and/or treating cancer, particularly prostate cancer. The proteins are useful as markers to indicate the presence or absence of cancer. This sequence represents a prostate cancer therapy associated CDNA. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly seconds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fanger GR,
Li SX, Wan
                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New fusion protein comprising prostate-specific polypeptides, or its immunogenic portions, useful for diagnosing, preventing and/or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Хu J,
                                                                                                                                                                                                                                                                 seqdata.uspto.gov/sequence.html?DocID=US20020192763.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 16; SEQ ID NO 471;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-352711/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mcneill
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(DILL/)
(MITC/)
(HARL/)
(JIAN/)
(KALO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RETT/)
(STOL/)
(DAYC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (VEDV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HOUG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MCNE/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HEND/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LISX/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FANG/
                                                           121
                                                                                    502
                                                                                                                                       562
        181
                                                                                                                                                                                       277;
                                                                                                            61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dillon DC, Mitcham JL, Harlocker SL, Jiang Y, GR, Retter MW, Stolk JA, Day CH, Vedvick TS, Wang A, Skeiky YAW, Hepler WT, Henderson RA, PD, Houghton RL, Y De Bassols CV, Foy TM;
                                                                                                                                                               1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HURAL J.

MCNEILL P D.

HOUGHTON R L.

Y DE BASSOLS C V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CARTER D.
LI S X.
WANG A.
WANG A.
SKEIKY Y A W.
HEPLER W T.
HENDERSON R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JIANG Y.
KALOS M D.
FANGER G R.
RETTER M W.
STOLK J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STOLK J A.
DAY C H.
VEDVICK T S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DILLON D C.
MITCHAM J L.
HARLOCKER S L.
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DILLON D
MITCHAM J
                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        particularly prostate cancer
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                                                 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCCATCTTGGG
AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAAC
                                                                                    TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                          TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAAGGACATTAGAAAATG
                                 AATTGATG
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                                                                                                                                                                                         Conservative
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Houghton RL,
                                                                                                                                                                                                                                         BP; 198 A; 189 C; 169
                                  95.7%;
99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             85pp; English.
                                                                                                                                                                                       0
                                                                                                                                                                                                  Score 266; DB 25;
Pred. No. 4.7e-76;
                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                         G; 256
                                                                                                                                                                                                                                         T; 0
                                                                                                                                                                                          0
                                                                                                                                                                                                                                          other;
                                                                                                                                                                                                                Length 812;
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Carter D;
Hural J;
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382

AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC

324

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RESULT 10
AAV62429
ID AAV62
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                                                                                                                                                                                                                                                          CC of the PCA3 gene. The PCA3 CDNA splice variant 3 sequence, isolated from a human primary prostatic tumour tissue cDNA library, CC was found in approximately 15% of the cDNA clones isolated. The CC was found in approximately 15% of the cDNA clones isolated. The CC invention claims for PCA3 CDNA variants and the proteins they encode. CC The invention also claims for antibodies against PCA3 protein. The CC antibodies are claimed to be useful for detecting PCA3 protein in CC immunoassay tests, for diagnosing, assessing and prognosing of prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin CC prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin CC redividual for treating PC, while determining elevated levels of PCA3 (as RNA or protein) is useful for detecting a predisposition CC odevelopment of PC, e.g. in prenatal tests. Detecting PCA3 protein CC allows differentiation between malignant and benign prostatic disease, and the level of PCA3 expression allows correlation with the grade of C tumour. PCA3 protein and its fragments are also claimed to be useful in vaccines for preventing PC; in drug screens for identifying specific (ant)agonists (potentially useful therapeutically) and for
                                                                                                                                        Query Match
Best Local S
Matches 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid encoding prostate cancer antigen prevention and treatment of prostatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-568347/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prostate cancer antigen (PCA3) cDNA splice variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV62429;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV62429 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the prostate cancer antigen (PCA3) cDNA splice variant 3 sequence comprising of exons 1, 3, and 4a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Pages 77-78; 111pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9845420-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prostate cancer antigen cDNA splice variant 3; PCA3; prostatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DIAG-) DIAGNOCURE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             323
   316
                                                                      256 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                       1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                          Similarity
                                                                                                                                                                                                                820
                      TCATCGATGAGCCTCGCCCTGTGCCCTGGTCCCGCTTGTGAGGGAAAGAACATTAGAAAATG
                                                                                                                                                                                                                                                 protein-DNA interactions
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TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                            BP; 262 A; 169 C; 191
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                                                                                                                                                         95.7%;
99.6%;
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                                                                                                                                      Score 266; DB 19;
Pred. No. 4.8e-76;
0; Mismatches 0
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                                                                                                                                                                                                              198 T;
                                                                                                                                                                           DB 19;
                                                                                                                                                                                                                0 other;
                                                                                                                                                                         Length 820;
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                                                                                                                                          Indels
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RESULT 11
AAD38815
ID AAD38
XX AAD38
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                                                                                                                                                                                                                   Query Match
Best Local S
Matches 277
                                                                                                                                                                                                                                                                                                                                                                                The invention relates to prostate specific proteins (PSP) and prostate specific nucleic acids (PSNA). Sequences of the invention are useful for identifying, diagnosing, monitoring, staging, imaging and treating prostate cancer and non-cancerous disease states in prostate. They are also useful for producing engineered prostate tissue for treatment and research. The PSNA sequences are used in gene therapy and for producing transgenic animals and cells. The invention is also used as vaccines. The present sequence is human PSNA cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated prostate specific polypeptide useful for identifying, diagnosing, monitoring, staging, imaging, and treating prostate cand non-cancerous disease states in prostate -
                                                                                                                                                                                                                                                                                                                              Sequence 876 BP; 275 A; 183 C; 207 G; 211 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; prostate specific protein; PSP; prostate specific nucleic acid; vaccine; transgenic; prostate cancer; gene therapy; transgenic animal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PSNA cDNA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 179; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-471216/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-2000; 2000US-233746P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-SEP-2001; 2001WO-US29386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PSNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-SEP-2002
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                                                                                                                                                                                                                                                   Local Similarity
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  345
                                                                                                                                                                                                                   277;
                                        61 TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                1 CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Recipon H, Cafferkey R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DIADEXUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TCATCGATGAGCCTCGCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                           CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGACGAGAAATCTTGATGGCTTCACAAGACATGCAAC 532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pro123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA;
                                                                                                                                                                                                                                                95.7%;
99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       876
                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                Score 266;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ali S;
                                                                                                                                                                                                                                                4.9e-76;
                                                                                                                                                                                                                                                                          DB 24;
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                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                В
                                                                                   CC CDNA splice variant 2 sequence comprising of exons 1, 3, 4a and CC cDNA splice variant 2 sequence comprising of exons 1, 3, 4a and CC 4b of the PCA3 gene. The PCA3 cDNA splice variant 2 sequence, do for the PCA3 gene. The PCA3 cDNA splice variant 2 sequence, cC isolated from a human primary prostatic tumour tissue cDNA library, cCC was found in approximately 65% of the cDNA clones isolated. The cCC invention claims for PCA3 cDNA variants and the proteins they encode. The invention also claims for antibodies against PCA3 protein. The CC minutunoassay tests, for diagnosing, assessing and prognosing of cCC prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin CC immunoassay tests, for diagnosing, assessing and prognosing of cCC prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin CC or radioisotope, and nucleic acids antisense to PCA3 cDNA are claimed cto be useful for treating PC, while determining elevated levels of CC pCA3 (as RNA or protein) is useful for detecting a predisposition cto development of PC, e.g. in prenatal tests. Detecting PCA3 protein CC allows differentiation between malignant and benign prostatic disease, cancer. PCA3 protein and its fragments are also claimed to be useful cin vaccines for preventing PC; in drug screens for identifying creating non-train-NNA interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid encoding prostate cancer antigen 3 prevention and treatment of prostatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Pages 76-77; lllpp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-568347/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bussemakers MJG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prostate cancer antigen (PCA3) cDNA splice variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV62428;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV62428 standard; cDNA; 1872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9845420-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DIAG-) DIAGNOCURE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121
                                                                     protein-DNA interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97US-0041836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-CA00346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antigen cDNA splice variant 2; PCA3; prostatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             240
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Query Match Best Local Similarity Matches 277; Conserv

Conservative

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Score 266; DB 1 Pred. No. 7e-76; 0; Mismatches

DB 19; 0

Length 1872; Indels

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Gaps

1,

95.7%;

1872

BP;

567 A; 389 C; 369 G; 539 T; 8

other;

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RESULT 13
AAV62427
ID AAV62
XX AAV62
XX AAV62
XX AAV62
XX Prost
XX
The present sequence represents the prostate cancer antigen (PCA3) cDNA splice variant 1 sequence comprising of exons 1, 2, 3, 4a and 4b of the PCA3 gene. The PCA3 CDNA splice variant 1 sequence, isolated from a human primary prostatic tumour tissue cDNA library, was found in approximately 5% of the cDNA clones isolated. The invention claims for PCA3 cDNA variants and the proteins they encode. The invention also claims for antibodies against PCA3 protein in the antibodies are claimed to be useful for detecting PCA3 protein in immunoassay tests, for diagnosing, assessing and prognosing of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prostate PC; ds.
                                                                                                                                                                                                                                                                                                                                                              New nucleic acid encoding prostate cancer antigen prevention and treatment of prostatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bussemakers MJG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV62427;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV62427 standard; cDNA; 2037
                                                                                                                                                                                                                                                                                                 Claim 3; Fig 2B-2J; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DIAG-) DIAGNOCURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1998-568347/48.
DB; AAW79736.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
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2019..2024
/*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen cDNA splice variant 1; PCA3; prostatic cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PCA3) cDNA splice variant 1.
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                                                                                                                                                                                                                                                                                                                                                                                             for diagnosis,
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RESULT 14
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XX AAA066
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XX Human;
KW Human;
KW immuno
XX Homo s
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Matches 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin or radioisotope, and nucleic acids antisense to PCA3 cDNA are claimed to be useful for treating PC, while determining elevated levels of PCA3 (as RNA or protein) is useful for detecting a predisposition to development of PC, e.g. in prenatal tests. Detecting PCA3 protein allows differentiation between malignant and benign prostatic disease, and the level of PCA3 expression allows correlation with the grade of tumour. PCA3 protein and its fragments are also claimed to be useful in vaccines for preventing PC, in drug screens for identifying specific (ant)agonists (potentially useful therapeutically) and for studying protein-DNA interactions.
                                                                                                      23-SEP-1998;
15-JAN-1999;
15-JAN-1999;
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                               Dillon
                                                                                                                                                                              14-JUL-1998;
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                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                                                                                                Human immunogenic prostate tumour protein cDNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA06688;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA06688 standard; cDNA;
   WPI; 2000-171268/15
                                                                                           09-APR-1999;
                                                                                                                                                     23-SEP-1998;
                                                                                                                                                                                                                                                                                                                                    immunogenic;
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                                                                                                                                                                                                                                                                                                                                  prostate cancer; diagnosis; tumour; gene therapy; detection;
genic; cytostatic; vaccine; ss.
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                                                             CORIXA
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                               Harlocker
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                                                                                      98US-0115453.

98US-0116134.

98US-0159812.

98US-0159822.

99US-0232149.

99US-0232880.

99US-0288946.
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Pred. No. 7.3e-76;
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                               Mitcham
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Claim 1; Page 260-261; 263pp; English.
                                                                                                      New polypeptide useful for treating comprises an immunogenic portion of
                                                                                                      and diagnosing prostate tumor
                                                                                                       prostate cancer
protein -
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The present invention describes isolated polypeptides, comprising an immunogenic portion of a prostate tumour protein (PTP). The polypeptides and polynucleotides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polypeptides and polynucleotides encoding them, antigen presenting cells which express the polypeptides, antibodies against the polypeptides and vaccines comprising them can be used for inhibiting the development of prostate cancer in a patient. The polypeptides can be used to generate antibodies or anti-idiotypic antibodies for passive immuno therapy. A portion of the polypucleotides encoding the polypeptides can be used as a probe or to modulate the expression of the polypeptides. AAA06241 to AAA06691 and AAV82000 to AAV82020 represent sequences used in the exemplification of the present invention. present invention.

Sequence 2229 BP; 654 A; 447 C; 481 G; 647 T; 0 other;

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                                                                                                                                              AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC
                                                       AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                                 TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                                                                                                                         CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCCGGCCGCCATCTTGGG
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AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 1341
                                         AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC
                                                                                     Conservative
                                                                                                                                                                                                                             95.7%;
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                                                                                                                                                                                                                             Score 266;
Pred. No. 7
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XX AASG38
XX Human |
XX Human |
XX Human |
XX Homo s
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09-MAY-2000;
12-MAY-2000;
    13-JUN-2000;
27-JUN-2000;
10-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human prostate cDNA sequence #431.
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                                                                                                                                                                                                                                                                                                                                              27-MAR-2001; 2001WO-US09919
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    ; 2000US-0536857.
; 2000US-0568100.
; 2000US-0570737.
; 2000US-0593793.
; 2000US-0605783.
; 2000US-0636215.
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Best Local Similarity
Matches 277; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                polynucleotides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fanger
Li SX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Χu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human prostate-specific polypeptides and polynucleotides useful the diagnosis and treatment of cancer, especially prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-AUG-2000; 2000US-0651236.

06-SEP-2000; 2000US-0657279.

02-OCT-2000; 2000US-0679426.

10-OCT-2000; 2000US-0685166.
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2229 BP; 654 A; 447 C; 481 G; 647 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to isolated prostate-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 385; 579pp; English.
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                                                                                                         AACGGGATTACAGATTTGAAATGAAGTCACCAAAGTGAGCATTACCAATGAGAGGAAAAC 240
                                                                                                                                                               TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG 120
                                                                                                                                                                                                                                                                                       CAGGAAGCACAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGCCGCCATCTTGGG
                                                                                                                                                                                                                                                                                                              CAGGAAGCACAAAAAGGAAGCACAGAGATCCCTGGGAGAAATGCCCGGGCCGCCATCTTGGG
                                 AGACGAGAAAATCTTGATGGCTTCACAAGACATGCAAC 278
                                                                                                                                              TCATCGATGAGCCTCGCCCTGTGCCTGGTCCCGCTTGTGAGGGAAGGACATTAGAAAATG
                                                                       AACGGGATTACAGATTTGAAATGAAGTCA-CAAAGTGAGCATTACCAATGAGAGGAAAAC 1379
                                                                                                                                                                                                                                                                                                                                                               Conservative
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99.6%;
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Carter D;
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Search completed: February Job time : 190 secs

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